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SEQUENCE LISTING

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TECH CENTER 1600/2900

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SUGANUMA, MASASHI
KAWABE, TAKUMI
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4212 + 1RT
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+210 + 339
· 2.13 · 11
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83
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· 210 / 340
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+ 212: PRT
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. 220>
\pm 223 > Description of Artificial Sequence: Synthetic
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Led Phe His Ser Fro Ser Ash Fro Glu Ash Led

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peptide
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·.12 · PRT
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Led Phe Ash Ser Pro Ser Blu The Blu Ash Led

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 - 312> PRT
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 <400> 1842
 Leu Tyr Gly Ser Pro Ser Met Pro Glu Asn Leu
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     peptide
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Leu Tyr Lys Ser Pro Ser Met Pro Glu Asn Leu
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peptide

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Leu Tyr Thr Ser Pro Ser Met Pro Glu Asn Leu
<210> 1852
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<210> 1853
<211> 11
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<400> 1853
Leu Tyr Tyr Ser Pro Ser Met Pro Glu Asn Leu
<210> 1854
<211> 11
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Leu Tyr Val Ser Pro Ser Met Pro Glu Asn Leu
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<211> 11
<212> PRT
<213> Artificial Sequence
<220>
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<212> PRT
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<400> 1856
Leu Tyr Arg Ser Pro Ser Arg Pro Glu Asn Leu
<210> 1857
<211> 11
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Leu Tyr Arg Ser Pro Ser Asn Pro Glu Asn Leu
                  5
<210> 1858
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Asp Pro Glu Asn Leu
<210> 1859
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<210> 1861
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<400> 1861
Leu Tyr Arg Ser Pro Ser Glu Pro Glu Asn Leu
<210> 1862
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<210> 1865
<211> 11
<212> PRT
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<400> 1865
Leu Tyr Arg Ser Pro Ser Leu Pro Glu Asn Leu
<210> 1866
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Lys Pro Glu Asn Leu
<210> 1867
4.2115 11
<212> PRT
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<400> 1867
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10.00

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<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
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<400> 1868
Leu Tyr Arg Ser Pro Ser Phe Pro Glu Asn Leu
<210> 1869
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<210> 1870
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Ser Pro Glu Asn Leu
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<400> 1871
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<400> 1873
Leu Tyr Arg Ser Pro Ser Tyr Pro Glu Asn Leu
<210> 1874
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<212> PRT
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<211> 11
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peptide

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<212> PRT
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<400> 1878
Leu Tyr Arg Ser Pro Ser Met Asp Glu Asn Leu
<210> 1879
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Met Cys Glu Asn Leu
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Leu Tyr Arg Ser Pro Ser Met Gln Glu Asn Leu
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<210> 1881
<211> 11
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<400> 1881
Leu Tyr Arg Ser Pro Ser Met Glu Glu Asn Leu
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<210> 1882
<211> 11
<212> PRT
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<211> 11
<212> PRT
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< 100> 1883
Leu Tyr Arg Ser Pro Ser Met His Glu Asn Leu
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<210> 1884
<211> 11
<212> PRT
<213> Artificial Sequence
. 220 -
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<211> 11
<212> PRT
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<211> 11
<212> PRT
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<400> 1886
Leu Tyr Arg Ser Pro Ser Met Lys Glu Asn Leu
<210> 1887
<211> 11
<212> PRT
<213> Artificial Sequence
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<400> 1887
Leu Tyr Arg Ser Pro Ser Met Met Glu Asn Leu
<210> 1888
<211> 11
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      peptide
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<210> 1889
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Met Pro Glu Asn Leu
<210> 1890
<211> 11
<212> PRT
<213> Artificial Sequence
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      peptide
<400> 1890
Leu Tyr Arg Ser Pro Ser Met Ser Glu Asn Leu
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                 5
<110> 1891
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Met Thr Glu Asn Leu
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<4(0> 1892
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<211> 11
<212> PRT
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<400> 1893
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                  5
<210> 1894
<211> 11
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 1894
Leu Tyr Arg Ser Pro Ser Met Val Glu Asn Leu
<210> 1895
<211> 9
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 1895
Tyr Gly Gly Pro Gly Gly Gly Asn
<210> 1896
<111> 11
<1112 > PRT
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<:400> 1896
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<223> Description of Artificial Sequence: Synthetic
     peptide
< 100> 1897
Leu Tyr Arg Ser Pro Ala Met Pro Glu Asn Leu
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<211> 8
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<223> Description of Artificial Sequence: Synthetic
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<400> 1898
Trp Tyr Arq Ser Pro Ser Tyr Tyr
<210> 1899
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
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<400> 1899
Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
<210> 1900
<211> 2055
<212> DNA
<113> Homo sapiens
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taastacete ettteeeeta gattteettt eattetgete aagtettege etgtgteega 120
tecetateta etttetetee tettgtagea ageeteagae teeaggettg agetaggttt 180
tgtttttctc ctggtgagaa ttcgaagacc atgtctacgg aactcttctc atccacaaga 240
gaggaaggaa getetggete aggaeceagt tttaggteta ateaaaggaa aatgttaaac 300
etgeteetgg agagagaeae tteetttace gtetgteeag atgteeetag aacteeagtg 360
qqcaaatttc ttggtgattc tgcaaaccta agcattttgt ctggaggaac cccaaaatgt 420
tgcctcgatc tttcgaatct tagcagtggg gagataactg ccactcaget taccacttct 480
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tcagatgaat taatggagtt ttccctgaaa gatcaagaag caaaggtgag cagaagtggc 840
ctatateget eccegtegat gecagagaae ttgaacagge caagactgaa geaggtggaa 900
aaattcaagg acaacacaat accagataaa gttaaaaaaa agtatttttc tggccaagga 960
aageteagga agggettatg titaaagaag acagtetete tgtgtgacat tactateact 1020
cagatgetgg aggaagatte taaccagggg cacetgattg gtgattttte caaggtatgt 1080
gegetgecaa eegtgteagg gaaacaccaa gatetgaagt atgteaacce agaaacagtg 1140
gctgccttac tgtcggggaa gttccagggt ctgattgaga agttttatgt cattgattgt 1200
cgctatccat atgagtatct gggaggacac atccagggag ccttaaactt atatagtcag 1260
gaagaactgt ttaacttett tetgaagaag eccategtee etttggacae ecagaagaga 1320
ataatcatcg tgttccactg tgaattctcc tcagagaggg gcccccgaat gtgccgctgt 1380
ctgcgtgaag aggacaggtc tctgaaccag tatcctgcat tgtactaccc agagctatat 1440
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atgageceat gataacatte cagecactgg etgetaacaa gteaceaaaa agacaetgea 1680
gaaaccctga gcagaaagag gccttctgga tggccaaacc caagattatt aaaagatgtc 1740
tetgeaaace aacaggetae caacttgtat ecaggeetgg gaatggatta ggttteagea 1800
gagetgaaag etggtggeag agteetggag etggetetat aaggeageet tgagttgeat 1860
agagatttgt attggttcag ggaactctgg cattcctttt cccaactcct catgtcttct 1920
cacaagccag ccaactettt etetetggge ttegggetat gcaagagegt tgtetacett 1980
etttetttgt atttteette tttgttteee eetettett ttttaaaaat ggaaaaataa 2040
acactacaga atgag
<210> 1901
<211> 472
<212> PRT
<213> Homo sapiens
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Met Ser Thr Glu Leu Phe Ser Ser Thr Arg Glu Glu Gly Ser Ser Gly
Ser Gly Pro Ser Phe Arg Ser Asn Gln Arg Lys Met Leu Asn Leu Leu
Leu Glu Arg Asp Thr Ser Phe Thr Val Cys Pro Asp Val Pro Arg Thr
Pro Val Gly Lys Phe Leu Gly Asp Ser Ala Asn Leu Ser Ile Leu Ser
```

Gly Gly Thr Pro Lys Cys Cys Leu Asp Leu Ser Asn Leu Ser Ser Gly 65 70 75 80

Glu Ile Thr Ala Thr Gln Leu Thr Tnr Ser Ala Asp Leu Asp Glu Thr 85 90 95

Gly His Leu Asp Ser Ser Leu Gln Glu Val His Leu Ala Gly Met Asn

100 105 110

His Asp Gln His Leu Met Lys Cys Ser Pro Ala Gln Leu Leu Cys Ser

His Asp Gln His Leu Met Lys Cys Ser Pro Ala Gln Leu Leu Cys Ser 115 - 120 - 125

the control of the co

| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu | Met | Lys | Tyr | Leu 165 | Gly | Ser | Pro | Ile | Thr 170 | Thr | Val | Pro | Lys | Leu 175 | Asp |
| Lys | Asn | Pro | Asn 180 | Leu | Gly | Glu | Asp | Gln 185 | Ala | Glu | Glu | Ile | Ser 190 | Asp | Glu |
| Leu | Met | Glu 195 | Phe | Ser | Leu | Lys | Asp 200 | Gln | Glu | Ala | Lys | Val 205 | Ser | Arg | Ser |
| Gly | Leu 210 | Tyr | Arg | Ser | Pro | Ser 215 | Met | Pro | Glu | Asn | Leu 220 | Asn | Arg | Pro | Arç |
| Leu 225 | Lys | Gln | Val | Glu | Lys 230 | Phe | Lys | Asp | Asn | Thr 235 | Ile | Pro | Asp | Lys | Val 240 |
| Lys | Lys | Lys | Tyr | Phe 245 | Ser | Gly | Gln | Gly | Lys 250 | Leu | Arg | Lys | Gly | Leu 255 | Cys |
| Leu | Lуs | Lys | Thr 260 | Val | Ser | Leu | Cys | Asp 265 | Ile | Thr | Ile | Thr | Gln 270 | Met | Let |
| Glu | Glu | Asp 275 | Ser | Asn | Gln | Gly | His 280 | Leu | Ile | Gly | Asp | Phe 285 | Ser | Lys | Val |
| Cys | Ala 290 | Leu | Pro | Thr | Val | Ser 295 | Gly | Lys | His | Gln | Asp 300 | Leu | Lys | Tyr | Val |
| Asn 305 | Pro | Glu | Thr | Val | Ala 310 | Ala | Leu | Leu | Ser | Gly 315 | Lys | Phe | Gln | Gly | Let 320 |
| Ile | Glu | Lys | Phe | Tyr 325 | Val | Ile | Asp | Cys | Arg 330 | Tyr | Pro | Tyr | Glu | Tyr 335 | Lei |
| Gly | Gly | His | 11e 340 | Gln | Gly | Ala | Leu | Asn 345 | Leu | Tyr | Ser | Gln | Glu 350 | Glu | Let |
| Phe | Asn | Phe 355 | Phe | Leu | Lys | Lys | Pro 360 | Ile | Val | Pro | Leu | Asp 365 | Thr | Gln | Lys |
| | Ile 370 | | Ile | | Phe | | Cys | | | Ser | | Glu | Arg | Gly | Pro |
| Arg 385 | Met | Cys | Arg | Cys | Leu 390 | Arg | Glu | Glu | Asp | Arg 395 | Ser | Leu | Asn | Gln | Ty: |
| Pro | Ala | Leu | Tyr | Tyr 405 | Pro | Glu | Leu | Tyr | Ile 410 | Leu | Lys | Gly | Gly | Tyr 415 | Arg |
| Asp | Phe | Phe | Pro 420 | Glu | Tyr | Met | Glu | Leu 425 | Cys | Glu | Pro | Gln | Ser 430 | Tyr | Cys |
| | | | | | - | | | | | | - | | - | · . | |

Leu Leu Val Lys Asp Met Ser Pro 465 470

<210> 1902

<211> 476

<212> PRT

<213> Homo sapiens

<400> 1902

Met Ala Val Pro Phe Val Glu Asp Trp Asp Leu Val Gln Thr Leu Gly

1 10 15

Glu Gly Ala Tyr Gly Glu Val Gln Leu Ala Val Asn Arg Val Thr Glu 20 25 30

Glu Ala Val Ala Val Lys Ile Val Asp Met Lys Arg Ala Val Asp Cys 35 40 45

Pro Glu Asn Ile Lys Lys Glu Ile Cys Ile Asn Lys Met Leu Asn His 50 55 60

Glu Asn Val Val Lys Phe Tyr Gly His Arg Arg Glu Gly Asn Ile Gln 65 70 75 80

Tyr Leu Phe Leu Glu Tyr Cys Ser Gly Gly Glu Leu Phe Asp Arg Ile 85 90 95

Glu Pro Asp Ile Gly Met Pro Glu Pro Asp Ala Gln Arg Phe Phe His
100 105 110

Gln Leu Met Ala Gly Val Val Tyr Leu His Gly Ile Gly Ile Thr His 115 120 125

Arg Asp Ile Lys Pro Glu Asn Leu Leu Asp Glu Arg Asp Asn Leu 130 135 140

Lys Ile Ser Asp Phe Gly Leu Ala Thr Val Phe Arg Tyr Asn Asn Arg 145 150 155 160

Glu Arg Leu Leu Asn Lys Met Cys Gly Thr Leu Pro Tyr Val Ala Pro \$165\$ \$170\$ \$175\$

Glu Leu Leu Lys Arg Glu Phe His Ala Glu Pro Val Asp Val Trp
180 185 190

Ser Cys Gly Ile Val Leu Thr Ala Met Leu Ala Gly Glu Leu Pro Trp 195 200 205

Asp Gln Pro Ser Asp Ser Cys Gln Glu Tyr Ser Asp Trp Lys Glu Lys 210 215 220

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Pro Asp Ile Lys Lys Asp Arg Trp Tyr Asn Lys Pro Leu Lys Lys Gly Ala Lys Arg Pro Arg Val Thr Ser Gly Gly Val Ser Glu Ser Pro Ser 280 Gly Phe Ser Lys His Ile Gln Ser Asn Leu Asp Phe Ser Pro Val Asn Ser Ala Ser Ser Glu Glu Asn Val Lys Tyr Ser Ser Ser Gln Pro Glu 310 315 Pro Arg Thr Gly Leu Ser Leu Trp Asp Thr Ser Pro Ser Tyr Ile Asp 325 330 Lys Leu Val Gln Gly Ile Ser Phe Ser Gln Pro Thr Cys Pro Asp His 345 Met Leu Asn Ser Gln Leu Leu Gly Thr Pro Gly Ser Ser Gln Asn 360 Pro Trp Gln Arg Leu Val Lys Arg Met Thr Arg Phe Phe Thr Lys Leu 375 Asp Ala Asp Lys Ser Tyr Gln Cys Leu Lys Glu Thr Cys Glu Lys Leu Gly Tyr Gln Trp Lys Lys Ser Cys Met Asn Gln Val Thr Ile Ser Thr 410 Thr Asp Arg Arg Asn Asn Lys Leu Ile Phe Lys Val Asn Leu Leu Glu

Met Asp Asp Lys Ile Leu Val Asp Phe Arg Leu Ser Lys Gly Asp Gly 435 440 445

Leu Glu Phe Lys Arg His Phe Leu Lys Ile Lys Gly Lys Leu Ile Asp
450 460

Ile Val Ser Ser Gln Lys Val Trp Leu Pro Ala Thr 465 470 475

<210> 1903

<211> 542

<212> PRT

<213> Homo sapiens

<400> 1903

Met Ser Arg Glu Ser Asp Val Glu Ala Gln Gln Ser His Gly Ser Ser 1 5 10 15

and the property of the property of the control of

- Ser Gln Ser Ser His Ser Ser Ser Gly Thr Leu Ser Ser Leu Glu Thr 50 55 60
- Val Ser Thr Gln Glu Leu Tyr Ser Ile Pro Glu Asp Gln Glu Pro Glu 65 70 75 80
- Asp Gln Glu Pro Glu Glu Pro Thr Pro Ala Pro Trp Ala Arg Leu Trp 85 90 95
- Ala Leu Gln Asp Gly Phe Ala Asn Leu Glu Cys Val Asn Asp Asn Tyr
 100 105 110
- Trp Phe Gly Arg Asp Lys Ser Cys Glu Tyr Cys Phe Asp Glu Pro Leu 115 120 125
- Leu Lys Arg Thr Asp Lys Tyr Arg Thr Tyr Ser Lys His Phe Arg 130 135 140
- Asp His Ser Gly Asn Gly Thr Phe Val Asn Thr Glu Leu Val Gly Lys 165 170 175
- Gly Lys Arg Arg Pro Leu Asn Asn Ser Glu Ile Ala Leu Ser Leu 180 185 190
- Ser Arg Asn Lys Val Phe Val Phe Phe Asp Leu Thr Val Asp Asp Gln 195 200 205
- Ser Val Tyr Pro Lys Ala Leu Arg Asp Glu Tyr Ile Met Ser Lys Thr 210 215 220
- Leu Gly Ser Gly Ala Cys Gly Glu Val Lys Leu Ala Phe Glu Arg Lys 225 230 235 240
- Thr Cys Lys Val Ala Ile Lys Ile Ile Ser Lys Arg Lys Phe Ala 245 250 255
- Ile Gly Ser Ala Arg Glu Ala Asp Pro Ala Leu Asn Val Glu Thr Glu 260 265 270
- Ile Glu Ile Leu Lys Lys Leu Asn His Pro Cys I.e Ile Lys Ile Lys 275 280 285
- Asn Phe Phe Asp Ala Glu Asp Tyr Tyr Ile Val Leu Glu Leu Met Glu 290 295 300
- Gly Glu Leu Phe Asp Lys Val Val Gly Asn Lys Arg Leu Lys Glu 305 310 315 320
- Ala Thr Cys Lys Leu Tyr Phe Tyr Gln Met Leu Leu Ala Val Gln Tyr

Leu Leu Ser Ser Gln Glu Glu Asp Cys Leu Ile Lys Ile Thr Asp Phe 355 360 365

Gly His Ser Lys Ile Leu Gly Glu Thr Ser Leu Met Arg Thr Leu Cys 370 375 380

Gly Thr Pro Thr Tyr Leu Ala Pro Glu Val Leu Val Ser Val Gly Thr 385 390 395 400

Ala Gly Tyr Asn Arg Ala Val Asp Cys Trp Ser Leu Gly Val Ile Leu 405 410 415

Phe Ile Cys Leu Ser Gly Tyr Pro Pro Phe Ser Glu His Arg Thr Gln 420 425 430

Val Ser Leu Lys Asp Gln Ile Thr Ser Gly Lys Tyr Asn Phe Ile Pro 435 440 445

Glu Val Trp Ala Glu Val Ser Glu Lys Ala Leu Asp Leu Val Lys Lys 450 455 460

Leu Leu Val Val Asp Pro Lys Ala Arg Phe Thr Thr Glu Glu Ala Leu 465 470 475 480

Arg His Pro Trp Leu Gln Asp Glu Asp Met Lys Arg Lys Phe Gln Asp 485 490 495

Leu Leu Ser Glu Glu Asn Glu Ser Thr Ala Leu Pro Gln Val Leu Ala 500 505 510

Gln Pro Ser Thr Ser Arg Lys Arg Pro Arg Glu Gly Glu Ala Glu Gly
515 520 525

Ala Glu Thr Thr Lys Arg Pro Ala Val Cys Ala Ala Val Leu 530 535 540

<210> 1904

<211> 12

<212> PRT

<213> Artificial Sequence

20005

<223> Description of Artificial Sequence: Synthetic peptide

<400> 1904

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<210> 1905

<211> 9

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peptide
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Gly Gly Arg Ser Pro Ala Met Pro Glu
<210> 1906
<211> 12
<212> PRT
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<210> 1907
<211> 11
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<220>
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<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1907
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<211> 11
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<220>
<221> MOD RES
<222: (8)
<223> Pro, Phe, Tyr or Trp
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<210> 1909
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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1909
Leu Tyr Thr Ser Pro Ser Tyr Xaa Glu Asn Leu
<210> 1910
<211> 11
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<223> Description of Artificial Sequence: Synthetic
      peptide
<220>
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<222> (8)
<223> Pro, Phe, Tyr or Trp
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<210> 1911
<211> 11
<212> PRT
<113> Artificial Sequence
<120>
<223> Description of Artificial Sequence: Synthetic
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
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<210> 1912
<211> 11
<212> PRT
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<211> 22
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<210> 1939
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Met Pro
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<210> 1944
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Gly Gly Asn
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<210> 1948
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